

1600

Does Not Comply Corrected Diskette Needed

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/392,822A

DATE: 09/11/2002 TIME: 09:06:09

Input Set : D:\Seqlist.txt

Output Set: N:\CRF4\09112002\I392822A.raw

ERRORED SEQUENCES

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60 <210> SEQ ID NO: 2
61 <211> LENGTH: 5836
62 <212> TYPE: DNA
63 <213> ORGANISM: H. sapiens
65 <400> SEQUENCE: 2
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67 atctgtgaca atattcacag tgtaatgcca tccagggaac tcaactgagc cttgatgtcc 120
68 agagattttt gtgttttttt ctqaqactqa qtctcqctct gtgccaqgct ggagtgcagt 180
69 ggtgcaacct tggctcactg caagctccgc ctcctgggtt cacgccattc tcctgcctca 240
70 gcctcctgag tagctgggac tacaggcacc cgccaccacg cctggctaat ttttttgtat 300
71 ttttagtaga gatggggttt cactgtgtta gccaggatgg tctcagtctc ctgacctcgt 360
72 gatetgeeca cettggeete ecaaagtget gggatgaeag gegtgageea eegegeetgg 420
73 ccgatateca gagatttttt ggggggetec ateaeacaga catgttgact gtetteatgg 480
74 ttgactttta gtatccagcc cctctagaaa tctagctgat atagtgtggc tcaaaacctt 540
75 cagcacaaat cacaccgtta gactatctgg tgtggcccaa accttcaggt gaacaaaggg 600
76 actotaatot ggcaggatat tocaaagcat tagagatgac otottgcaaa gaaaaagaaa 660
78 gaggggaaac gcctgaggtc tttgagcaag gtcagtcctc tgttgcacag tctccctcac 780
79 agggteattg tgacgateaa atgtggteae gtgtatgagg caccageaea tgeetggete 840
80 tggggagtgc cgtgtaagtg tatgcttgca ctgctgaatg cttgggatgt gtcagggatt 900
81 atcttcagca cttacagatg ctcatctcat cctcacagca tcactatggg atgggtatta 960
82 ctggcctcat ttgatggaga aagtggctgt ggctcagaaa ggggggacca ctagaccagg 1020
83 gacactotgg atgotgggga otocagagao catgaccact caccaactgo agagaaatta 1080
84 attgtggcct gatgtccctg tcctggagag ggtggaggtg gaccttcact aacctcctac 1140
85 cttgaccete tettttaggg etetttetga eetecaceat ggtactagga eeceattgta 1200
86 ttctgtaccc tcttgactct atgaccccca ctgcccactg catccagctg ggtcccctcc 1260
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136 attettgaga cacagetegg getggateaa gtgagetaet etggttttat tgaacagetg 4260
     137 aaataaccaa ctttttggaa attgatgaaa tcttacggag ttaacagtgg aggtaccagg 4320
     138 getettaaga gtteeegatt etettetgag aetacaaatt gtgattttge atgeeacett 4380
     139 aatotttttt ttttttttt taaatogagg tttcagtctc attctatttc ccaggctgga 4440
     140 gttcaatage gtgatcacag ctcactgtag cettgaacte etggeettaa gagattetee 4500
     141 tgcttcggtc tcccaatagc taagactaca gtagtccacc accatatcca gataattttt 4560
     142 aaattttttg gggggccggg cacagtggct cacgcctgta atcccaacac catgggaggc 4620
     143 tgagatgggt ggatcacgag gtcaggagtt tgagaccagc ctgaccaaca tggtgaaact 4680
     144 ctytetetae taaaaaaaaa aaaaatagaa aaattageeg ggegtggtgg cacaeggeae 4740
     145 ctgtaatccc agctactgag gaggctgagg caggagaatc acttgaaccc agaaggcaga 4800
     146 ggttgcaatg agccgagatt gcgccactgc actccagcct gggtgacaga gtgagactct 4860
     147 gtctcaaaaa aaaaaaattt ttttttttt tttgtagaga tggatcttgc tttgtttctc 4920
     148 tggttggcct tgaactectg getteaagtg atcetectae ettggeeteg gaaagtgttg 4980
     149 ggattacagg cgtgagccac catgactgac ctgtcgttaa tcttgaggta cataaacctg 5040
     150 gctcctaaag gctaaaggct aaatatttgt tggagaaggg gcattggatt ttgcatgagg 5100
     151 atgattetga eetgggaggg eaggteagea ggeatetetg ttgcaeagat agagtgtaea 5160
     152 ggtctggaga acaaggagtg gggggttatt ggaattccac attgtttgct gcacgttgga 5220
     153 ttttgaaatg ctagggaact ttgggagact catatttctg ggctagagga tctgtggacc 5280
     154 acaagatett tttatgatga cagtageaat gtatetgtgg agetggatte tgggttggga 5340
     155 gtgcaaggaa aagaatgtac taaatgccaa gacatctatt tcaggagcat gaggaataaa 5400
     156 agttetagtt tetggtetea gagtggtgea gggateaggg agteteaeaa teteetgagt 5460
     157 gctggtgtct tagggcacac tgggtcttgg agtgcaaagg atctaggcac gtgaggcttt 5520
     158 gtatgaagaa teggggateg tacceaecee etgtttetgt tteateetgg geatgtetee 5580
     159 totgoottig toccotagat gaagtotoca tgagotacaa gggcotggtg catocagggt 5640
     160 gatctagtaa ttgcagaaca gcaagtgcta gctctccctc cccttccaca gctctgggtg 5700
     161 tgggaggggg ttgtccagcc tccagcagca tggggagggc cttggtcagc ctctgggtgc 5760
     162 cagcagggca ggggcggagt cctggggaat gaaggtttta tagggctcct gggggaggct 5820/
E--> 163 ccccaqcccc aagctt
                                                                           5836 (210)
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     165 <212> TYPE: DNA
     166 <213> ORGANISM: H. sapiens
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E--> 168 <400> SEQUENCE: 3
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     171 cagggetece agtggeetga gaaccattgg acccaggace cattacttet agggtaagga 180
     172 aggtacaaac accagatcca accatggtct ggggggacag ctgtcaaatg cctaaaaata 240
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     174 ggcaacccaa agtccaaatc caggtgagca ggtgcaccaa atgcccagag atatgacgag 360
     175 gcaagaagtg aaggaaccac ccctgcatca aatgttttgc atgggaagga gaagggggtt 420
     176 gctcatgttc ccaatccagg agaatgcatt tgggatctgc cttcttctca ctccttggtt 480
     177 agcaagacta agcaaccagg actctggatt tgggggaaaga cgtttatttg tggaggccag 540
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    180 tetgttgata teagatggee ceattttetg tacetteaca gaaggacaca ggetagggte 720
     181 tgtgcatggc cttcatcccc ggggccatgt gaggacagca ggtgggaaag atcatgggtc 780
    182 ctcctgggtc ctgcagggcc agaacattca tcacccatac tgacctccta gatgggaatg 840
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183 gcttccctgg ggctgggcca acggggcctg ggcaggggag aaaggacgtc aggggacagg 900 184 gaggaagggt catcgagacc cagcctggaa ggttcttgtc tctgaccatc caggatttac 960 RAW SEQUENCE LISTING DATE: 09/11/2002 PATENT APPLICATION: US/09/392,822A TIME: 09:06:09

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381 acctecetea geegggeeea eageeeagea gggteeacee teaceegggt caccteggee 12780
     382 cacqtcctcc tegeceteeg agetecteae aeggaetetg teageteete cetgeageet 12840
     383 ateggeegee cacetgagge ttgteggeeg eccaettgag geetgtegge tgeeetetge 12900
     384 aggcagetee tgteecetae acceeeteet teeceggget cagetgaaag ggcgteteee 12960
     385 agggcagete cetgtgatet ceaggacage teagtetete acaggeteeg acgeeeceta 13020
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     394 cctcttgccc tccagggggt gacattgcac acagacatca ctcaggaaac ggattcccct 13560
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     397 accetgttet gecaggggte cetgtgeaac tecagaetee etectggtae caccatgggg 13740
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     406 agacccaagg acagaacaca gcagaggtca gcactgggga agacaggttg tcctcccagg 14280
     407 ggatgggggt ccatccacct tgccgaaaag atttgtctga ggaactgaaa atagaaggga 14340
     408 aaaaagagga gggacaaaag aggcagaaat gagaggggag gggacagagg acacctgaat 14400
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                                                                                 , in sort hand return
     415 gacaggagtc acaccagaaa aatcaaattg aactggaatt ggaaaggggc aggaaaacct 14820
     416 caagagttot attttootag ttaattgtoa otggooacta ogtttttaaa aatoataata 14880
     417 actgcatcag atgacacttt aaataaaaac ataaccaggg catgaaacac tgtcctcatc 14940
     418 cgcctaccgc ggacattgga aaataagccc caggctgtgg agggccctgg gaaccctcat 15000
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     427 aagatgtetg aacacccaac acatagcact ggagatatga getegacaag agttteteag 180
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429 attgcgctca cgattggctt gaagaagcaa ggactgtggg aggtgggctt tgtagtaaca 300

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     618 gacaggagaa teggttgagt etgggagtte aaggetacag ggagetgega teaegeeget 11640
     619 gcactccage ctgggaaaca gagtgagact gtctcagaat ttttttaaaa aagaatcagt 11700
     620 gatcatccca acccctgttg ctgttcatcc tgagcctgcc ttctctggct ttgttcccta 11760
     621 gatcacatct ccatgatcca taggecetge ccaatctgae etcacacegt gggaatgeet 11820
                                                                                  , most hand return
     622 ccagactgat ctagtatgtg tggaacagca agtgctggct ctccctcccc ttccacagct 11880
     623 ctgggtgtgg gagggggttg tccagcctcc agcagcatgg ggagggcctt ggtcagcatc 11940
     624 taggtgccaa cagggcaagg gcggggtcct ggagaatgaa ggctttatag ggctcctcag 12000 🗸
E--> 625 ggaggccccc cagccccaaa ctgcaccacc tggccgtgga caccggt
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626 <211> LENGTH: 858 627 <212> TYPE: DNA 628 <213> ORGANISM: H. sapiens W--> 630 <210> SEQ ID NO: E--> 630 <400> SEQUENCE: 5 631 cgagcggccc ctcagcttcg gcgcccagcc ccgcaaggct cccggtgacc actagagggc 60 632 qqqaqqaqct cctgqccaqt qqtqqaqaqt qqcaaqqaaq gaccctaggq ttcatcgqaq 120 633 cccaggttta ctcccttaag tggaaatttc ttcccccact cctccttggc tttctccaag 180 634 gaqqqaaccc aggctqctqq aaaqtccqqc tgggqcqqqq actqtqgqtt cagggqagaa 240 635 cggggtgtgg aacgggacag ggagcggtta gaagggtggg gctattccgg gaagtggtgg 300 636 qqqqaqqqaq cccaaaacta qcacctagtc cactcattat ccagccctct tatttctcgg 360 637 ccgctctgct tcagtggacc cggggagggc ggggaagtgg agtgggagac ctaggggtgg 420 638 getteeegae ettgetgtae aggaeetega eetagetgge tttgtteeee ateeceaegt 480 639 tagttgttgc cctgaggcta aaactagagc ccaggggccc caagttccag actgcccctc 540 640 cccctcccc cggagccagg gagtggttgg tgaaaggggg aggccagctg gagaacaaac 600 641 gggtagtcag ggggttgagc gattagagcc cttgtaccct acccaggaat ggttggggag 660 642 gaggaggaag aggtaggagg taggggaggg ggcggggttt tgtcacctgt cacctgctcg 720 643 ctgtgcctag ggcgggcggg cggggagtgg ggggaccggt ataaagcggt aggcgcctgt 780 644 georgetica ecteteaage agecagegee tgeetgaate tgttetgeee ecteeceaee 840 645 catttcacca ccaccatg 858 RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/11/2002 PATENT APPLICATION: US/09/392,822A TIME: 09:06:10

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Output Set: N:\CRF4\09112002\I392822A.raw

Skipped Sequences(NEW RULES):

Sequence(s)_missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number

<400> sequence id number 000

Seq#:3,4,5

VERIFICATION SUMMARY DATE: 09/11/2002

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Input Set : D:\Seqlist.txt

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- L:163 M:254 E: No. of Bases conflict, LENGTH:Input:3 Counted:5838 SEQ:2
- L:163 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9
- $\texttt{L:}163 \texttt{ M:}252 \texttt{ E: No. of Seq. differs, <211> \texttt{LENGTH:}Input:}5836 \texttt{ Found:}5838 \texttt{ SEQ:}2$
- $L:168\ M:282\ W:$ Numeric Field Identifier Missing, <210> is required.
- L:168 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:3
- M:254 Repeated in SeqNo=2
- L:419 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
- L:419 M:252 E: No. of Seq. differs, <211> LENGTH:Input:15056 Found:15058 SEQ:0
- L:424 M:282 W: Numeric Field Identifier Missing, <210> is required.
- L:424 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:4
- L:625 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
- L:625 M:252 E: No. of Seq. differs, <211> LENGTH:Input:12047 Found:12049 SEQ:0
- L:630 M:282 W: Numeric Field Identifier Missing, <210> is required.
- L:630 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:5
- L:647 M:216 E: (34) Seq. #s missing, SEQ ID NOS: 3 thru 5